SEQUENCE LISTING

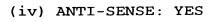
(1) GENERAL INFORMATION:

- (i) APPLICANT: Agrawal, Sudhir
- (ii) TITLE OF INVENTION: NOVEL HIV-SPECIFIC SYNTHETIC OLIGONUCLEOTIDES AND METHODS OF THEIR USE
- (iii) NUMBER OF SEQUENCES: 6
 - (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: Hale and Dorr LLP
 - (B) STREET: 60 State Street
 - (C) CITY: Boston
 - (D) STATE: Massachusetts
 - (E) COUNTRY: USA
 - (F) ZIP: 02109
 - (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Floppy disk
 - (B) COMPUTER: IBM PC compatible
 - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 - (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
- (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER:
 - (B) FILING DATE:
 - (C) CLASSIFICATION:
- (vii) PRIOR APPLICATION DATA:
 - (A) APPLICATION NUMBER: US
 - (B) FILING DATE:
- (viii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: Kerner, Ann-Louise
 - (B) REGISTRATION NUMBER: 33,523
 - (C) REFERENCE/DOCKET NUMBER: HYZ-069
 - (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: 617-526-6000
 - (B) TELEFAX: 617-526-5000
- (2) INFORMATION FOR SEQ ID NO:1:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 21 base pairs
 - (B) TYPE: nucleic acid

	(C) STRANDEDNESS: single (D) TOPOLOGY: linear		
(ii)	MOLECULE TYPE: DNA/RNA		
(iii)	HYPOTHETICAL: NO		
(iv)	ANTI-SENSE: YES		
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:1:		
UCGCACCCA	AT CTCTCCCUU C	21	
(2) INFOR	MATION FOR SEQ ID NO:2:		
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 21 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear		
(ii)	MOLECULE TYPE: DNA		
(iii)	HYPOTHETICAL: NO		
(iv)	ANTI-SENSE: YES		
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:2:		
TCGCACCCA	T CTCTCTCCTT C		21
(2) INFOR	MATION FOR SEQ ID NO:3:		
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 21 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear		
(ii)	MOLECULE TYPE: DNA/RNA		
(iii)	HYPOTHETICAL: NO		
(iv)	ANTI-SENSE: YES		
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:3:		
CGCACCCAT	ר יירייריירינווור וו	21	

(2) INFORMATION FOR SEQ ID NO:4:		
(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 21 base pairs(B) TYPE: nucleic acid(C) STRANDEDNESS: single(D) TOPOLOGY: linear		
(ii) MOLECULE TYPE: DNA		
(iii) HYPOTHETICAL: NO		
(iv) ANTI-SENSE: YES		
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:		
CGCACCCATC TCTCTCCTTC T		21
(2) INFORMATION FOR SEQ ID NO:5:		
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 22 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 		
(ii) MOLECULE TYPE: DNA		
(iii) HYPOTHETICAL: NO		
(iv) ANTI-SENSE: NO ,		
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:		
TCGCACCCAT CTCTCTCTT CT	22	
(2) INORMATION FOR SEQ ID NO:6:		
(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 21 base pairs(B) TYPE: nucleic acid(C) STRANDEDNESS: single(D) TOPOLOGY: linear	·	
(ii) MOLECULE TYPE: DNA/RNA		

(iii) HYPOTHETICAL: NO



(xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:

TCGCACCCAT CTCTCTCTT C

21